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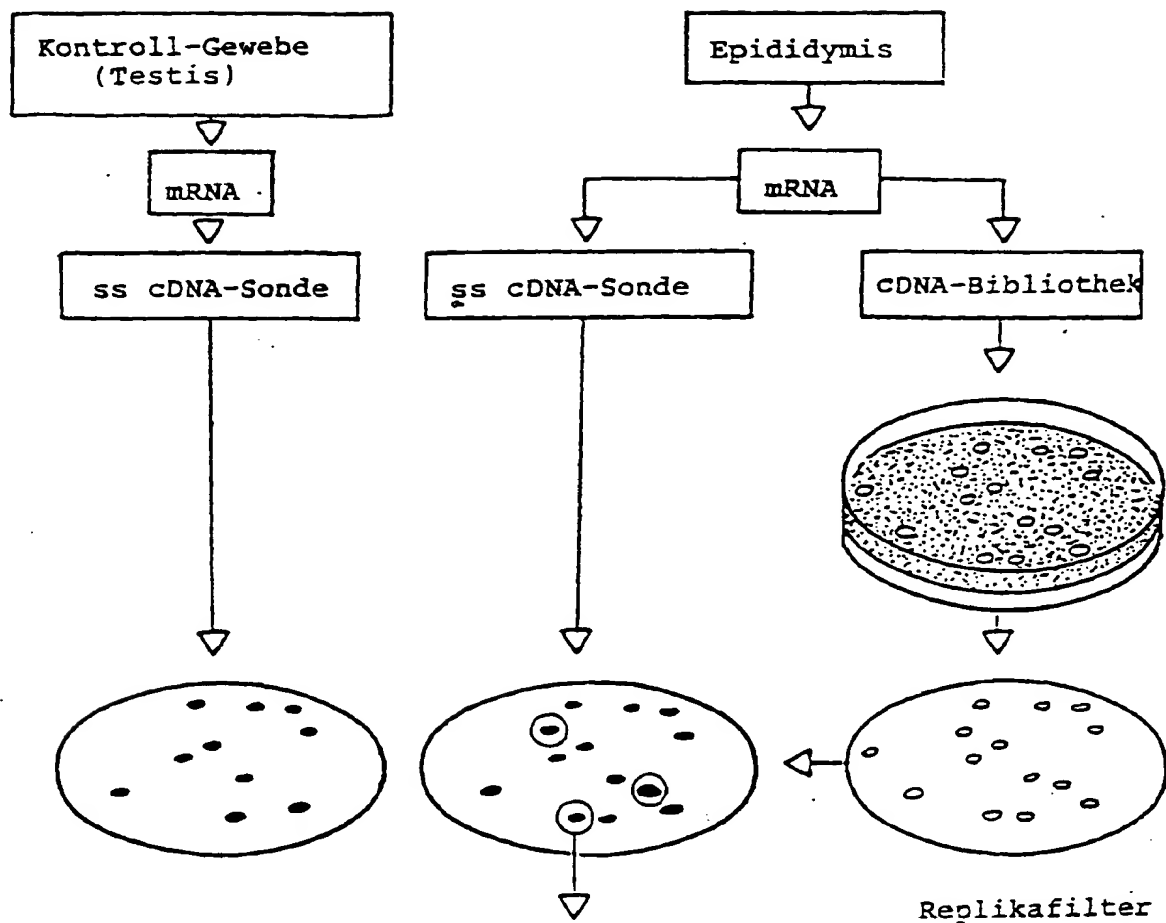
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Figur 1

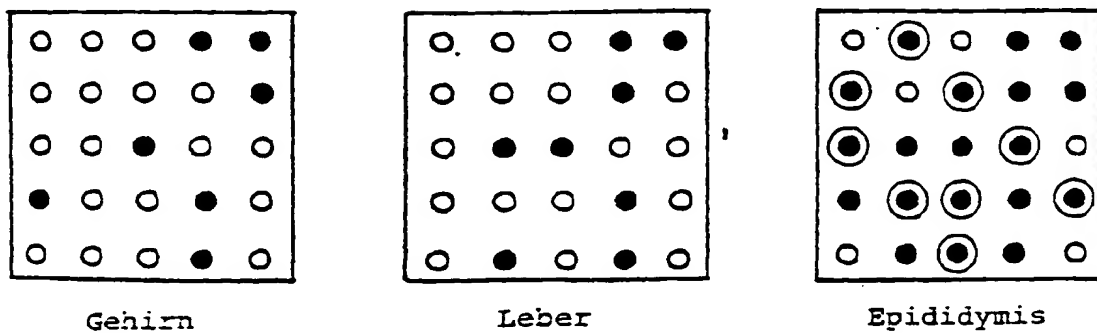
a)

## PRIMÄR-SCREENING



b)

## SEKUNDÄR-SCREENING



Figur 2

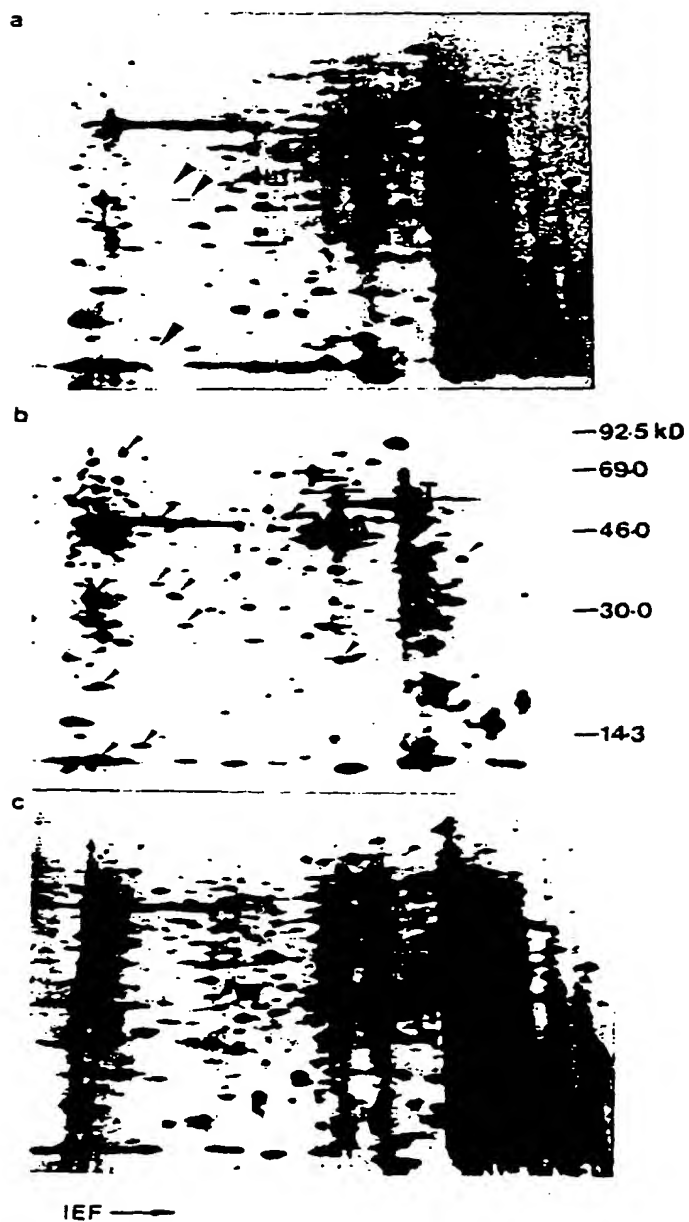
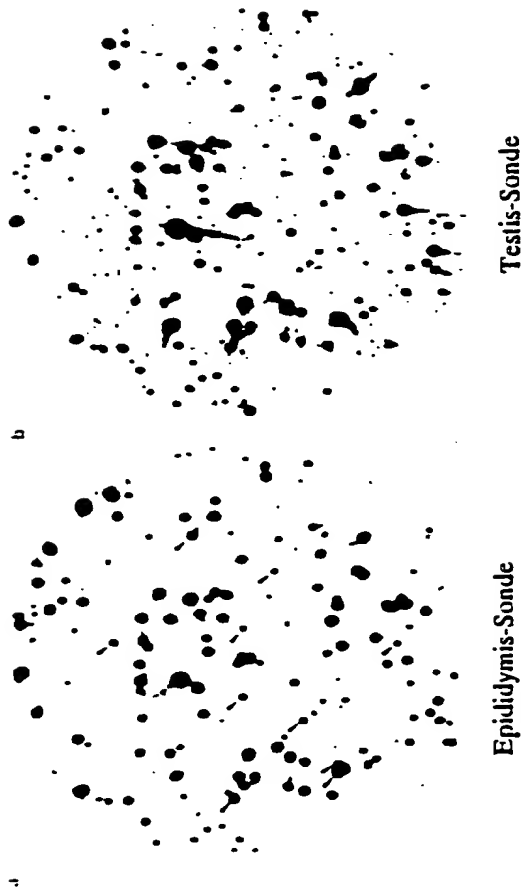


Figure 3



Figur 4

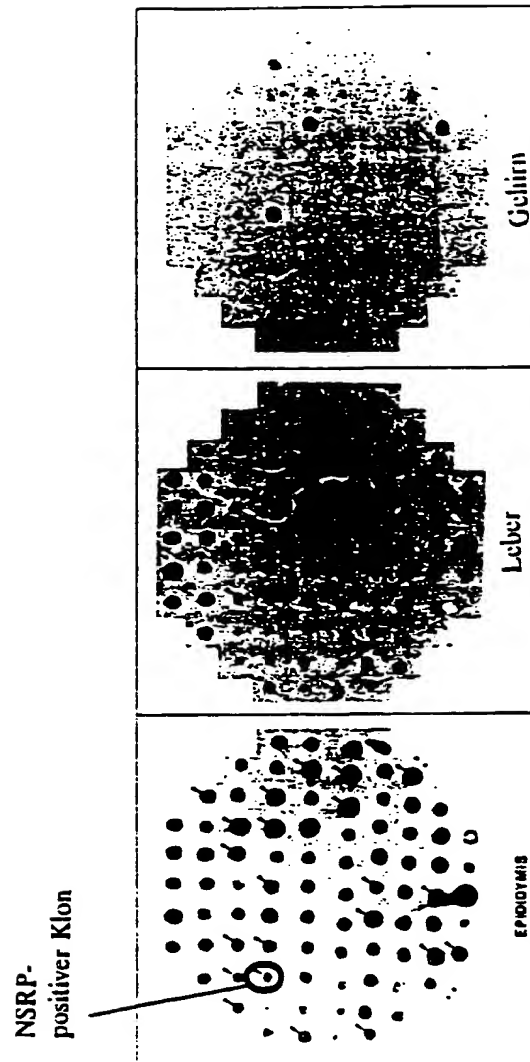


Figure 5

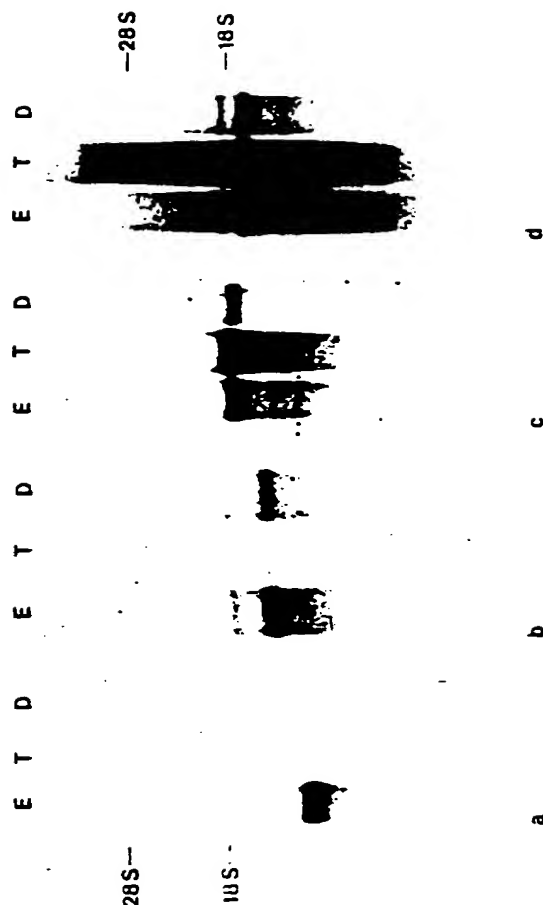
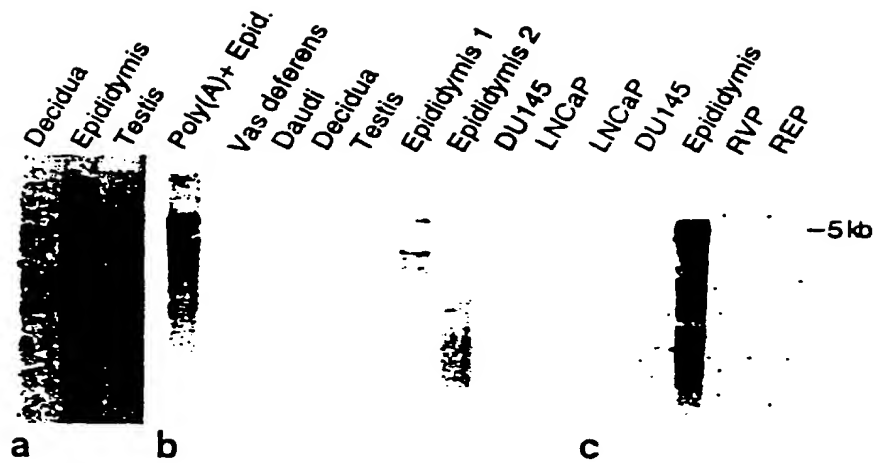
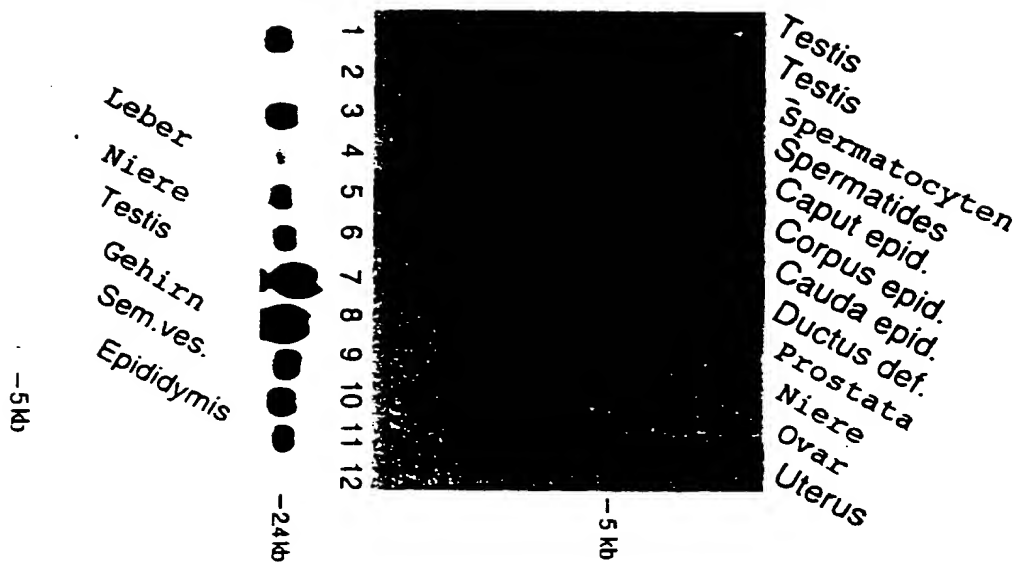


Figure 6

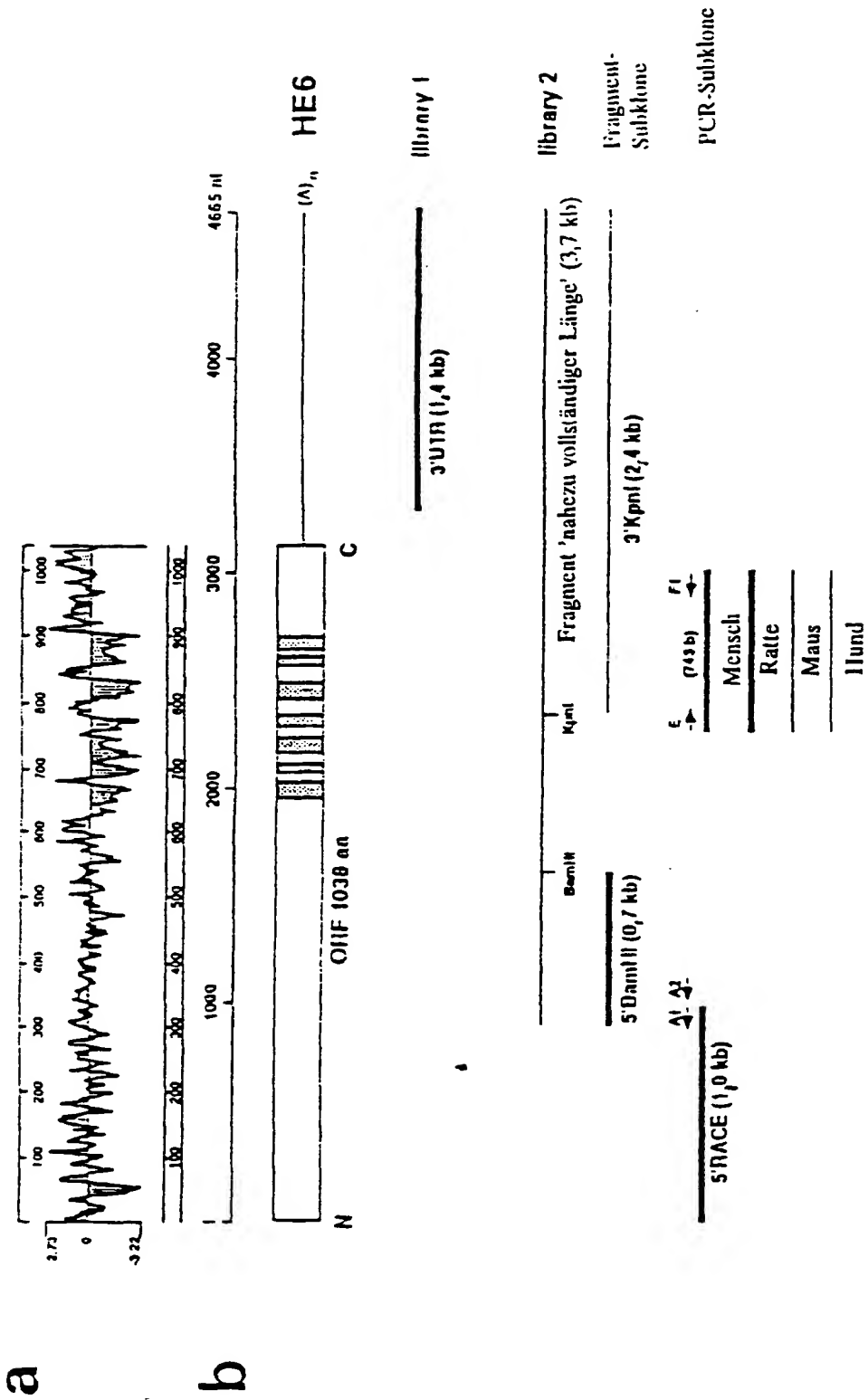


Figur 7





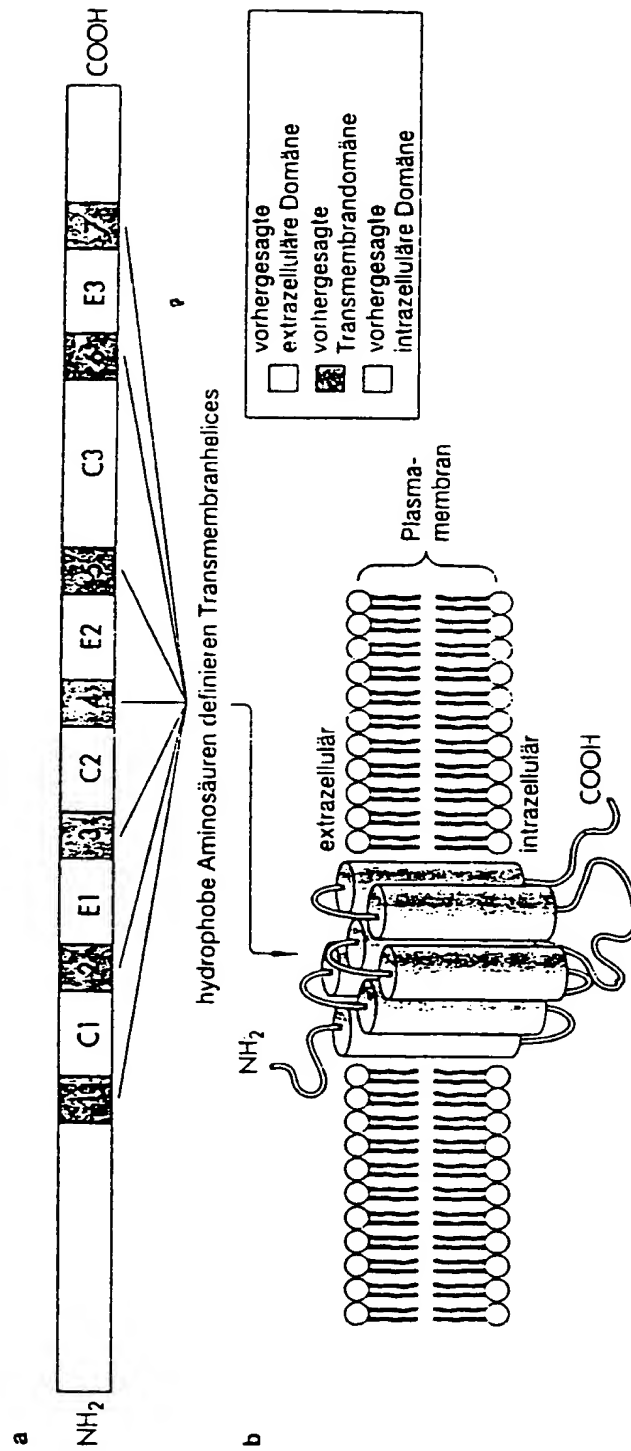
Figur 8a, b



Fragmente, die als Sonden verwendet werden, sind als schwarze Balken hervorgehoben.

Figur 8c

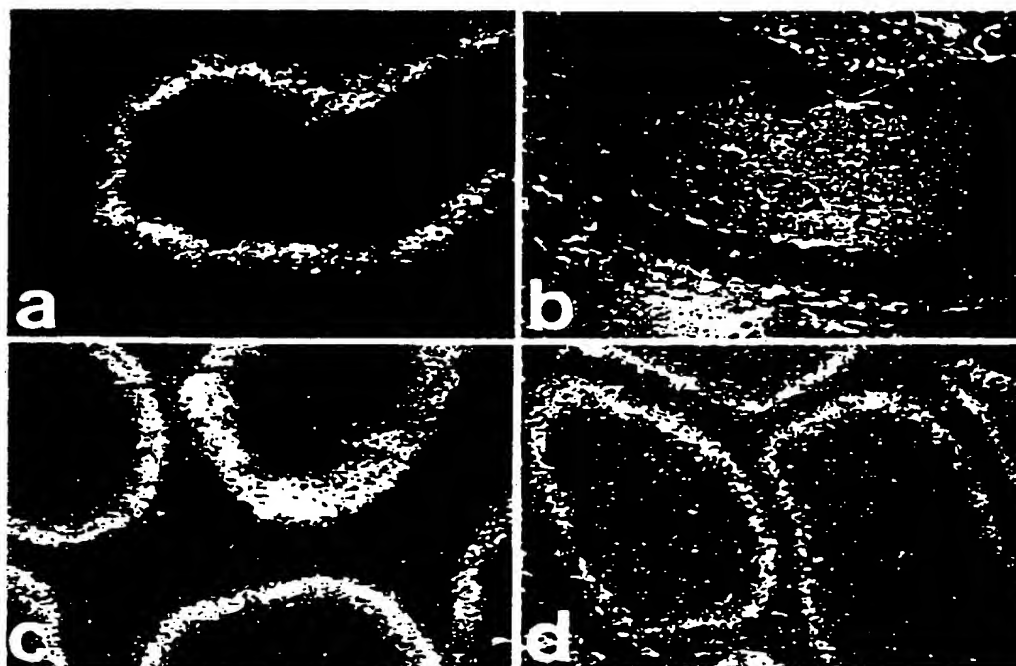
## 17. Signaltransfer durch Membranen



durch Schleifen verbunden, die abwechselnd außerhalb (E) und innerhalb (C) der Zelle liegen. Der N-Terminus der Rezeptoren weist nach außen, der C-Terminus dagegen ins Zellinnere. (b) Modell der mutmaßlichen Struktur des gefalteten Rezeptors.

17.3 Aufbau von Rezeptoren mit sieben Helices. (a) Lineare Darstellung des Rezeptorproteins. Sieben Blöcke aus hydrophoben Aminosäuren (1-7) nehmen jeweils die Form einer Helix an, die sich vermutlich durch die Membran erstreckt. Die Helices sind

Figur 9



Line 34 : BNSDOC tag : No converter for FO FP Output Format PS CC EP KC PA

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